2 tor	CASTrora Corrected by the STIC Sy. The Branch CRE Processing Date: 4/5/200/
UM Sorkal	Number: 09/405, 2/6
$\Box$ .	Changed a me members as a second seco
	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings of subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence jumbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:
<u> </u>	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Doloted extra, invalid, hoadings used by an applicant, specifically:
	Deleted: non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as
	Insorted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted
	Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (error due to a Patentin bug). Sequences corrected:
	Other:
	James T.

Action. DO NOT send a copy of this form.